

gina

FILE 'HOME' ENTERED AT 10:40:23 ON 13 NOV 2003

=> file reg

COST IN U.S. DOLLARS

SINCE FILE

TOTAL

ENTRY

SESSION

FULL ESTIMATED COST

0.21

0.21

FILE 'REGISTRY' ENTERED AT 10:40:32 ON 13 NOV 2003

USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.

PLEASE SEE "HELP USAGETERMS" FOR DETAILS.

COPYRIGHT (C) 2003 American Chemical Society (ACS)

Property values tagged with IC are from the ZIC/VINITI data file provided by InfoChem.

STRUCTURE FILE UPDATES: 12 NOV 2003 HIGHEST RN 616193-58-9

DICTIONARY FILE UPDATES: 12 NOV 2003 HIGHEST RN 616193-58-9

TSCA INFORMATION NOW CURRENT THROUGH JULY 14, 2003

Please note that search-term pricing does apply when conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. See HELP PROPERTIES for more information. See STN Note 27, Searching Properties in the CAS Registry File, for complete details:

<http://www.cas.org/ONLINE/STN/STNOTES/stnotes27.pdf>

=>

Uploading diamide.str

L1 STRUCTURE UPLOADED

=> d

L1 HAS NO ANSWERS

L1 STR

*** STRUCTURE DIAGRAM IS NOT AVAILABLE ***

Structure attributes must be viewed using STN Express query preparation.

=> s l1 sam

SAMPLE SEARCH INITIATED 10:40:59 FILE 'REGISTRY'

SAMPLE SCREEN SEARCH COMPLETED - 257 TO ITERATE

100.0% PROCESSED 257 ITERATIONS

0 ANSWERS

SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE **COMPLETE**

BATCH **COMPLETE**

PROJECTED ITERATIONS: 4179 TO 6101

PROJECTED ANSWERS: 0 TO 0

L2 0 SEA SSS SAM L1

=> s l1 full

FULL SEARCH INITIATED 10:41:05 FILE 'REGISTRY'

FULL SCREEN SEARCH COMPLETED - 5200 TO ITERATE

gina

100.0% PROCESSED 5200 ITERATIONS
SEARCH TIME: 00.00.03

0 ANSWERS

L3 0 SEA SSS FUL L1

=> logoff

ALL L# QUERIES AND ANSWER SETS ARE DELETED AT LOGOFF

LOGOFF? (Y)/N/HOLD:y

COST IN U.S. DOLLARS

SINCE FILE

TOTAL

ENTRY

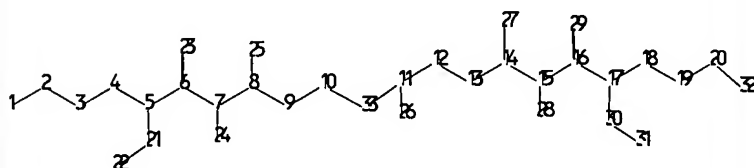
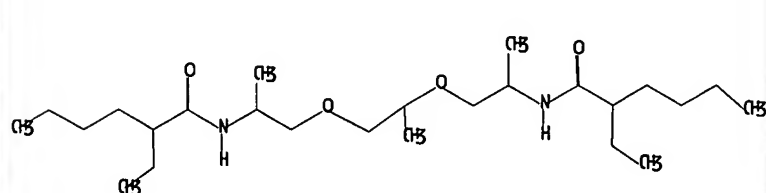
SESSION

FULL ESTIMATED COST

148.15

148.36

STN INTERNATIONAL LOGOFF AT 10:41:23 ON 13 NOV 2003



chain nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25
26 27 28 29 30 31 32 33

chain bonds :

1-2 2-3 3-4 4-5 5-6 5-21 6-7 6-23 7-8 7-24 8-9 8-25 9-10 10-33 11-26 11-12
11-33 12-13 13-14 14-15 14-27 15-16 15-28 16-17 16-29 17-18 17-30 18-19 19-20
20-32 21-22 30-31

exact/norm bonds :

6-7 6-23 7-8 9-10 10-33 11-12 12-13 14-15 15-16 16-29

exact bonds :

1-2 2-3 3-4 4-5 5-6 5-21 7-24 8-9 8-25 11-26 11-33 13-14 14-27 15-28 16-17
17-18 17-30 18-19 19-20 20-32 21-22 30-31

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:CLASS 9:CLASS 10:CLASS
11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS
20:CLASS 21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS
29:CLASS 30:CLASS 31:CLASS 32:CLASS 33:CLASS